C. KAUFMAN

1646

#29

RAW SEQUENCE LISTING DATE: 12/12/2000 PATENT APPLICATION: US/09/157,289E TIME: 12:52:44

Input Set : A:\P1134R2 REVISED.txt
Output Set: N:\CRF3\12122000\I157289E.raw

```
3 <110> APPLICANT: ASHKENAZI, AVI J
         BOTSTEIN, DAVID
         DODGE, KELLY H.
 6
         GURNEY, AUSTIN L.
         KIM, KYUNG JIN
 8
         LAWRENCE, DAVID A.
 9
         PITTI, ROBERT
10
         ROY, MARGARET A
11
         TUMAS', DANJEL B
12
         WOOD, WILLIAM I.
14 <120> TITLE OF INVENTION: DCR3 Polypeptide, A TNFR Homolog
16 <130> FILE REFERENCE: P1134R2 REVISED
18 <140> CURRENT APPLICATION NUMBER: US 09/157,289E
19 <141> CURRENT FILING DATE: 1998-09-18
21 <150> PRIOR APPLICATION NUMBER: US 60/059,288
22 <151> PRIOR FILING DATE: 1997-09-18
24 <150> PRIOR APPLICATION NUMBER: US 60/094,640
25 <151> PRIOR FILING DATE: 1998-07-30
27 <160> NUMBER OF SEQ ID NOS: 18
29 <210> SEO ID NO: 1
30 <211> LENGTH: 300
31 <212> TYPE: PRT
32 <213> ORGANISM: Homo sapiens
34 <400> SEQUENCE: 1
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36
     1
                      5
38
    Leu Ala Leu Pro Ala Leu Leu Pro Val Pro Ala Val Arg Gly Val
39
                     20
                                          25
4.1
    Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu
42
                     35
44
    Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg
45
                     50
                                          55
47
    Pro Cys Arg Arg Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro
48
    Arg His Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr
51
                     8.0
                                          85
53
    Cys Asn Val Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys
54
                     95
                                         100
56
    His Ala Thr His Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe
57
                    110
                                                             120
                                         115
59
    Ala His Ala Gly Phe Cys Leu Glu His Ala Ser Cys Pro Pro Gly
60
                    125
                                         130
    Ala Gly Val Ile Ala Pro Gly Thr Pro Ser Gln Asn Thr Gln Cys
63
                    1.40
                                         145
                                                             150
65
   Gln Pro Cys Pro Pro Gly Thr Phe Ser Ala Ser Ser Ser Ser
                    155
                                         160
   Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala Leu Gly Leu Ala
```

## ENTERED

## RECEIVED

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Input Set : A:\P1134R2 REVISED.txt
Output Set: N:\CRF3\12122000\I157289E.raw

```
Leu Asn Val Pro Gly Ser Ser Ser His Asp Thr Leu Cys Thr Ser
     71
                         185
                                                                  195
     72
                                              190
     74
         Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala Glu Glu
     75
                         200
                                              205
     77
         Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile Ser
     78
                         215
                                              220
                                                                  225
     80
        The Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu Ala Pro Glu
     81.
                         230
                                              235
        Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu
     83
     84
                         245
                                              250
                                                                  255
        Lys Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly
     86
     87
                         260
                                              265
     89
        Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met
     90
                         275
                                             280
                                                                  285
     92
        Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His
     93
                         290
                                              295
     95 <210> SEO TD NO: 2
     96 <211> LENGTH: 1114
     97 <212> TYPE: DNA
     98 <213> ORGANISM: Homo sapiens
     100 <220> FEATURE:
     101 <221> NAME/KEY: Unsure
     102 <222> LOCATION: 1090
    103 <223> OTHER INFORMATION: Unknown base
    105 <400> SEQUENCE: 2
    106 tecgcaggeg gaceggggge aaaggaggtg geatgteggt eaggeacage 50
         agggteetgt gteegegetg ageegegete teeetgetee ageaaggaee 100
    110
         atgagggege tggagggee aggeetgteg etgetgtgee tggtgttgge 150
    11.2
         getgeetgee etgetgeegg tgeeggetgt acgeggagtg geagaaacae 200
         ccacctacce etggegggac geagagaeag gggagegget ggtgtgegee 250
     1.1.4
         cagtyccccc caggeacett tgtgcagegg cegtgcegec gagacagece 300
         cacgacgtgt gycccgtgtc caccgcgcca ctacacgcag ttctggaact 350
    118
    120
         acctggageg etgeogetac tgcaacgtce tetgegggga gegtgaggag 400
    122
         gaggeaeggg ettgecaege cacecaeaac egtgeetgee getgeegeae 450
         cygettette gegeaegety gtttetgett ggageaegea tegtgteeac 500
         ctggtgccgg cgtgattgcc ccgggcaccc ccagccagaa cacgcagtgc 550
    126
         cagoogtgoo coccaggoae ettotoagoe agoagotoca gotoagagoa 600
    128
    130
         gtgccagccc cacegcaact gcaeggecet gggcctggcc ctcaatgtgc 650
         caggetette eteccatgae accetgtgea ecagetgeae tggetteece 700
         ctcagcacca gggtaccagg agctgaggag 'tgtgagcgtg ccgtcatcga 750
    134
    136
         ctttgtggct ttccaggaca tctccatcaa gaggctgcag cggctgctgc 800
    138
         aggecetega ggeceeggag ggetggggte egaeaceaag ggegggeege 850
         geggeettge agetgaaget gegteggegg etcaeggage teetggggge 900
         gcaggacggg gcgctgctgg tgcggctgct gcaggcgctg cgcgtggcca 950
    142
    144
         ggatgeeegg getggagegg agegteegtg agegetteet eeetgtgeae 1000
    146
         tgatectgge eccetettat ttattetaca teettggeae eccaettgea 1050
         ctgaaagagg cttttttta aatagaagaa atgaggtttn ttaaaaaaaa 1100
W--> 148
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## RECEIVED

DEC 2 7 2000

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aaaaaaaaa aaaa 1114

150

Input Set : A:\P1134R2 REVISED.txt
Output Set: N:\CRF3\12122000\1157289E.raw

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152 <210> SEQ ID NO: 3
     153 <211> LENGTH: 491
      154 <212> TYPE: DNA
      155 <213> ORGANISM: Unknown
      157 <220> FEATURE:
      158 <223> OTHER INFORMATION: Unknown organism
      160 <220> FEATURE:
     161 <221> NAME/KEY: unsure / 162 <222> LOCATION: 62, 73, 86, 98
      163 <223> OTHER INFORMATION: unknown base
      165 <400> SEQUENCE: 3
      166 geogagacag ceccaegacg tgtggecegt gtecaecgeg ceaetacacg 50
W--> 168 cagttetgga antaactgga genetgeege tactgnaacg_teetetgngg 100
      170 ggagcgtgag gaggaggcac gggettgcca cgccacccac aaccgtgeet 150
      172 geogetyceg caccygette tregegeacy crygthrety criggayeae 200
      174
           geategtgte cacetggtge eggegtgatt geecegggea ecceeageea 250
           gaacacgeag tgcctagecg tgccccceag geacettete agceageage 300
     178 tecageteag ageagtgeea geoceaeege aactgeaegg eeetgggeet. 350
      180 gyccotcaat gtgccaggot ottootcoca tgacaccotg tgcaccagot 400
      182 geactygett ecceetcage accagggtae caggagetga ggagtgtgag 450
     184 cgtgccgtca tegactttgt ggctttccag gacatetcca t 491
     186 <210> SEQ ID NO: 4
     187 <211> LENGTH: 73
     188 <212> TYPE: DNA
     189 <213> ORGANISM: Unknown
     191 <220> FEATURE:
     192 <223> OTHER INFORMATION: Unknown organism
     194 <400> SEQUENCE: 4
     195 geogagacag ceccaegacg tgtggeeegt gtecaeegeg ceaetacaeg 50
     197 cattetggaa etaeetggag ege 73
     1.99 <210> SEQ ID NO: 5
     200 <211> LENGTH: 271
     201 <212> TYPE: DNA
     202 <213> ORGANISM: Unknown
     204 <220> FEATURE:
     205 <223> OTHER INFORMATION: Unknown organism
     207 <220> FEATURE:
     208 <221> NAME/KEY: unsure/
209 <222> LOCATION: 42, 62, 73, 86, 98, 106, 120, 122, 153, 167, 184, 220, 233
     210 <223> OTHER INFORMATION: unknown base
     212 <400> SEQUENCE: 5
W--> 213 gccgagacag ccccacgacg tgtggcccgt gtccaccgcg cnactacacg 50
W--> 215 <u>cagttctgga</u> a<u>ntaactgga</u> <u>genetgeege</u> <u>tactgnaacg</u> <u>tectetgngg</u> 100
W--> 217 <u>ggagenfgag</u> <u>gaggaggean</u> <u>gngettgeea</u> <u>cgecac</u>cac <u>aaccgegeet</u> 150
W--> 219 gcngctgcag caccggnttc ttcgcgcacg ctgnttctg cttggagcac 200
W--> 221 gcatcgtgtc cacctggtgn cggcgtgatt gcnccgggca cccccagcca 250
     223 gaacacgcat gcaaagccgt g 271
     225 <210> SEQ LD NO: 6
     226 <211> LENGTH: 201
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Input Set : A:\P1134R2 REVISED.txt
Output Set: N:\CRF3\12122000\1157289E.raw

227 <212> TYPE: DNA 228 <213> ORGANISM: Unknown 230 <220> FEATURE: 231 <223> OTHER INFORMATION: Unknown organism 233 <220> FEATURE: 234 <221> NAME/KEY: unsure 235 <222> LOCATION: 182 236 <223> OTHER INFORMATION: unknown base 238 <400> SEQUENCE: 6 239 geagttetgg aactacetgg agegetgeeg etactgeaac gteetetgeg 50 241 gggagcgtga ggaggaggca egggettgec aegccaccca eaaccgtgec 100 243 tgccgctqcc gcaccqgctt cttcgcgcac gcgcgtttct gcttggagca 150 W--> 245 cgcatcgtgt ccacctggtg ccggcgtgat thecccgggc acccccagec 200 247 a 201 249 <210> SEQ ID NO: 7 250 <211> LENGTH: 277 251 <212> TYPE: DNA 252 <213> ORGANISM: Unknown 254 <220> FEATURE: 255 <223> OTHER INFORMATION: Unknown organism 257 <220> FEATURE: 258 <221> NAME/KEY: unsure 259 <222> LOCATION: 142 260 <223> OTHER INFORMATION: unknown base 262 <400> SEQUENCE: 7 263 gaggggcccc caggagtggt ggccggaggt gtggcagggg tcaggttgct 50 265 ggtcccagcc ttgcaccctg agctaggaca ccagttcccc tgaccctgtt 100 W--> 267 cttccctcct <u>ggctgcaggc ac</u>ccccagcc agaacacgca gnccagccgt 150 269 gcccccagg cacettetea gccagcaget ccageteaga geagtgccag 200 271 ecceacegea actgcaegge cetgggeetg geecteaatg tgccaggete 250 273 ttcctcccat gacaccetgt gcaccag 277 275 <210> SEQ ID NO: 8 276 <211> LENGTH: 199 277 <212> TYPE: DNA 278 <213> ORGANISM: Unknown 280 <220> FEATURE: 281 <223> OTHER INFORMATION: Unknown organism 283 <400> SEQUENCE: 8 284 gcatcgtgtc cacctggtgc cggcgtgatt gccccgggca cccccagcca 50 286 gaacacgcag gcctagccgt gccccccagg caccttetca gccagcagct 100 288 ccageteaga geagtgeeag ceceaecgea actgeaegge eetgggeetg 150 290 gccctcaatg tgccaggete tteeteecat gacaccetgt gcaccaget 199 292 <210> SEQ ID NO: 9 293 <211> LENGTH: 226 294 <212> TYPE: DNA 295 <213> ORGANISM: Unknown 297 <220> FEATURE:

298 <223> OTHER INFORMATION: Unknown organism

300 <220> FEATURE:

Input Set : A:\P1134R2 REVISED.txt
Output Set: N:\CRF3\12122000\I157289E.raw

301 <221> NAME/KEY: unsure -302 <222> LOCATION: 4, 9, 12, 165 303 <223> OTHER INFORMATION: unknown base 305 <400> SEQUENCE: 9 W--> 306 agengtgene enéaggeace tteteageca geagtteeag eteagageag 50 308 Egccaycccc accycaacty cacygootty gycotygoot toaatgtgcc 100 310 aggetettee teecatgaca egetgtgeac cagetgeact ggetteecec 150 W--> 312 tcagcaccag ggtancagga gctgaggagt gtgagcgtgc cgtcatcgac 200 314 tttgtggctt tccaggacat ctccat 226 316 <210> SEQ ID NO: 10 317 <211> LENGTH: 283 318 <212> TYPE: DNA 319 <213> ORGANISM: Homo sapiens 321 <220> FEATURE: 322 <221> NAME/KEY: Unsure 323 <222> LOCATION: 1-283 324 <223> OTHER INFORMATION: Unknown organism 326 <220> FEATURE: 327 <221> NAME/KEY: unsure 328 <222> LOCATION: 27, 64, 140 329 <223> OTHER INFORMATION: unknown base 331 <400> SEQUENCE: 10 W--> 332 cttgtccacc tggtgccggc gtgattnccc gggcaccccc agccagaaca 50 W--> 334 cgcagtgcca gccntcccc caggcactt ctcagccagc agctccagct 100
W--> 336 cagagcagtg ccagcccae cgcaactgca acgccctgn ctggccctca 150
338 atgtgccagg ctcttcctcc catgacacc tgtgcaccag ctgcactggc 200 340 theecectea geaccagggt accaggaget gaggagtgtg agegtgeegt 250 342 categaettt gtggetttee aggaeatete cat 283 344 <210> SEQ ID NO: 11 345 <211> LENGTH: 21 346 <212> TYPE: DNA 347 <213> ORGANISM: Unknown 349 <220> FEATURE: 350 <223> OTHER INFORMATION: Unknown organism 352 <400> SEQUENCE: 11 353 cacgctggtt tctgcttgga g 21 355 <210> SEQ ID NO: 12 356 <211> LENGTH: 22 357 <212> TYPE: DNA 358 <213> ORGANISM: Unknown 360 <220> FEATURE: 361 <223> OTHER INFORMATION: Unknown organism 363 <400> SEQUENCE: 12 364 agetggtgea cagggtgtea tg 22 366 <210> SEQ 1D NO: 13 367 <211> LENGTH: 53 368 <212> TYPE: DNA 369 <213> ORGANISM: Unknown

371 <220> FEATURE:

VERIFICATION SUMMARY

DATE: 12/12/2000 TIME: 12:52:45

PATENT APPLICATION: US/09/157,289E

Input Set : A:\P1134R2 REVISED.txt
Output Set: N:\CRF3\12122000\I157289E.raw

L:148 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:213 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:215 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:217 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:219 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:221 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:221 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:306 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:312 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:332 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:334 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:336 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10